

The open-source Treatment Planning Toolkit “matRad”

Introduction & Update on ongoing Developments and related Projects

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Treatment Planning

- computerized process
- dose is numerically simulated and optimized

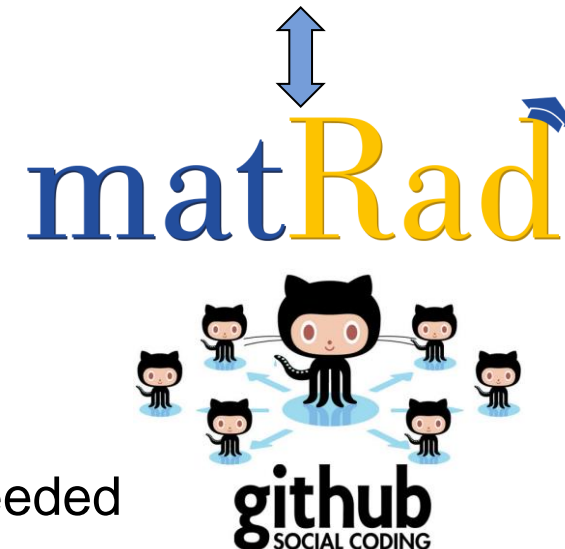
Commercial solutions are closed systems (Black Box)



Research needs flexible, accessible software

Examples for research topics:

- Biological Optimization (RBE, effect, mixed-modality)
 - Probabilistic dose calculation & optimization
- low-level access to dose calculation / optimization needed



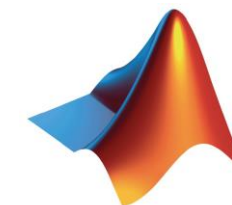
What is **matRad**?

- toolkit for three-dimensional intensity-modulated treatment planning for photons, protons and carbon ions
- Entirely written in Matlab & open source
- matRad implements well-established radiotherapy algorithms for **research & education**

Properties:

- **open-source code, patients and machine files on GitHub**
- **graphical user interface**
- **Non-linear constrained dose optimization (IPOPT)**
- **Import & export functionalities (DICOM, binary formats)**
- **No Matlab? → Octave compatibility & downloadable standalone**

Why? Supporting open science, reproducibility and education



www.matrad.org

Team



dkfz.

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IN THE HELMHOLTZ ASSOCIATION



DFG

Deutsche
Forschungsgemeinschaft

BA 2279/3-1
WA 4707/1-1

DKFZ Development team

Niklas Wahl

Lucas Burigo

Amit Ben Antony Bennan

Noa Homolka

Oliver Jäkel

HIT cooperation

Benjamin Ackermann

Swantje Ecker

Malte Ellerbrock

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Advisors

Martin Siggel

Peter Ziegenhein

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Hans-Peter Wieser

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Thomas Klinge

Verena Böswald

Henning Mescher

Alexander Stadler

Guiseppe Pezzano

Lucas-Raphael Müller

Hubert Gabrys

Silke Ulrich

Oliver Schrenk

Paul Meder

Other Contributors

Eric Crhistiansen (Carleton University)

Steven van de Water (PSI)

H.-P. Wieser *et al.* (all authors above in **bold**), “Development of the open-source dose calculation and optimization toolkit matRad,” *Med Phys*, vol. 44, no. 6, pp. 2556–2568, 2017, doi: [10.1002/mp.12251](https://doi.org/10.1002/mp.12251).

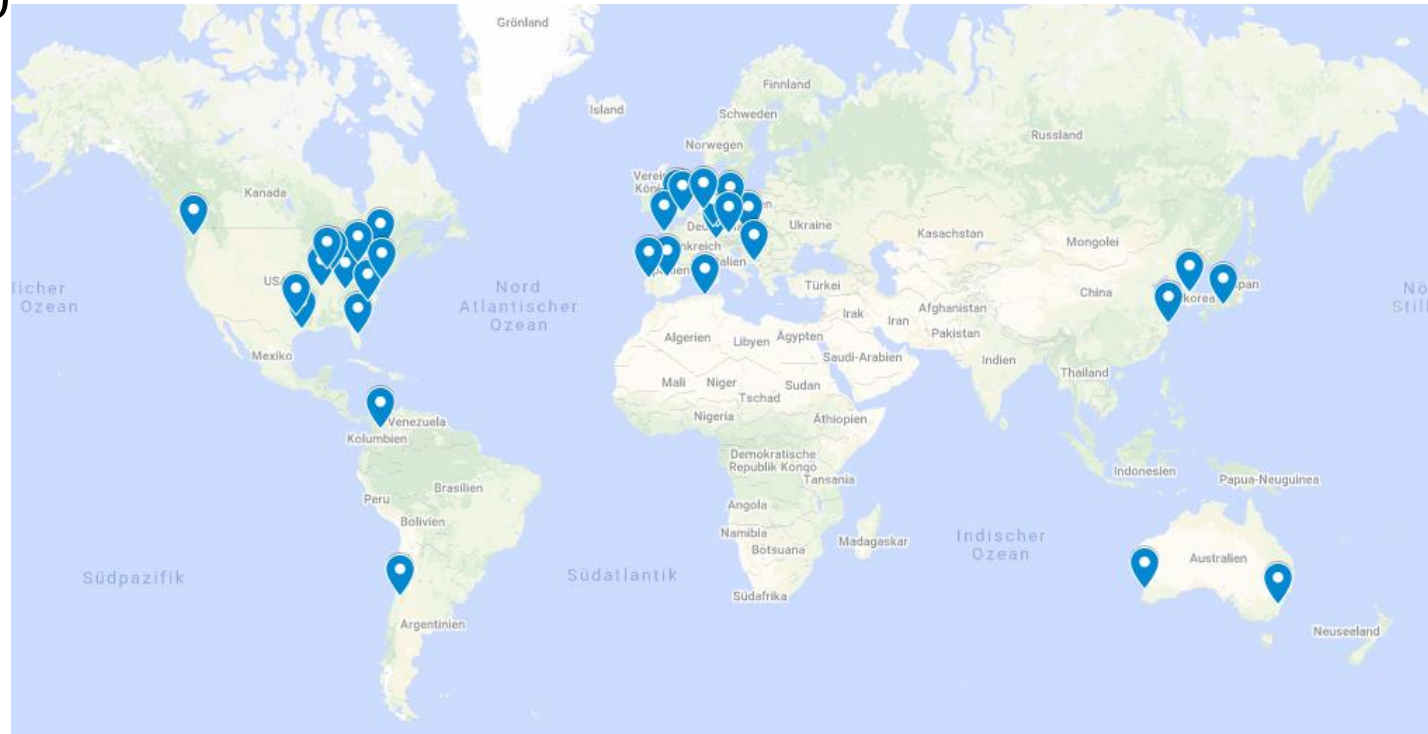
All Code Contributors: <https://github.com/e0404/matRad/blob/master/AUTHORS.txt>

More than 25 confirmed institutes somehow working with matRad

Start: 22. Januar 2015

1st commit 12cdfc1

- Github-Forks: >110



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University of Zurich UZH



UNIVERSITÄT HEIDELBERG ZUKUNFT SEIT 1386



THE UNIVERSITY OF WESTERN AUSTRALIA

More than 25 confirmed institutes somehow working with matRad

• Github

e0404 / matRad

Unwatch 26 Unstar 123 Fork 118

Code Issues 18 Pull requests 5 Discussions Actions Projects 3 Wiki Security Insights Settings

Filters is:issue is:open Labels 10 Milestones 1 New issue

18 Open 183 Closed

Author Label Projects Milestones Assignee Sort

- [Feature Request] Separate dose threshold for matRad_gammaIndex enhancement #515 opened 21 days ago by wahl
- Question about VMAT branches bug question 41 #499 opened on 31 Mar by chh105
- question about the dij structure enhancement question 18 #477 opened on 11 Jan by chh105
- [BUG] Color-Scale in GUI wrong after re-optimization with different number of fractions bug #472 opened on 28 Dec 2020 by wahl
- Sensible structure set resampling for optimization? enhancement question #462 opened on 6 Nov 2020 by wahl
- matRad has issues with 2D patients bug enhancement #324 opened on 23 Jan 2019 by wahl
- Continue GUI development with MATLAB guide or switch to code? enhancement help wanted question 4 #294 opened on 19 Sep 2018 by wahl
- DICOM import of Raystation PBS plans enhancement help wanted 4 #280 opened on 19 Jul 2018 by sebastianuber

2015




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Development of the open-source dose calculation and optimization toolkit matRad

Hans-Peter Wieser , Eduardo Cisternas, Niklas Wahl, Silke Ulrich, Alexander Stadler, Henning Mescher, Lucas-Raphael Müller, Thomas Klinge, Hubert Gabrys, Lucas Burigo, Andrea Mairani, Swantje Ecker, Benjamin Ackermann, Malte Ellerbrock, Katia Parodi, Oliver Jäkel, Mark Bangert

*Wieser et al., 2017, Med Phys 44(6)
among top 20 downloaded Med Phys
papers in 2017*

- 3D dose calculation (validated)

Photons: SVD pencil-beam algorithm + sequencing
MC interface to ompMC (open source)

Protons: Pencil-Beam algorithm + const. RBE
MC interface to MCsquare (open source)

Carbon ions: Pencil-Beam algorithm + biol. effect / RBE

- Base data

Patient data (CORT data set) & DICOM Import

Physical (& biological) base data for photon LINAC as well as a proton and a carbon machine

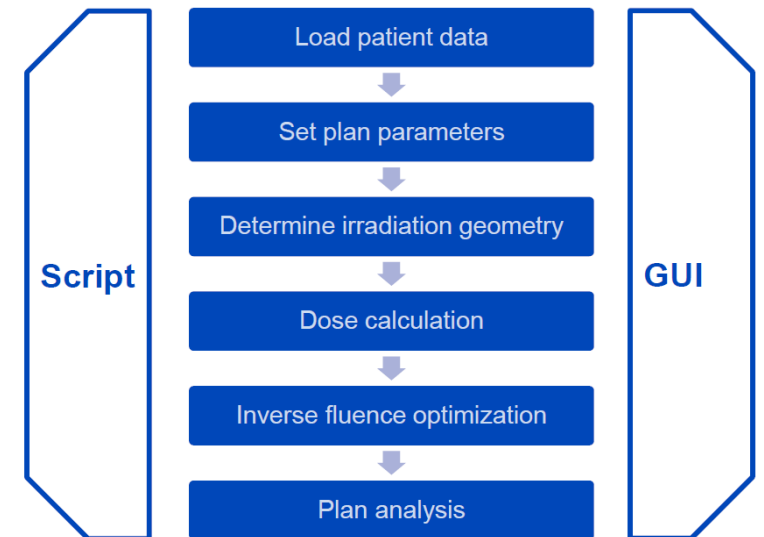
- Inverse planning with new optimization interface

Photons: Physical dose optimization & DAO

Protons: + Constant RBE optimization

Carbon-ions: + RBE (1.1 or variable) or effect optimization

- Scripting & Graphical User Interface



Workflow

Refresh | Load *.mat data | Calc. influence Mx | Optimize | Save to GUI
 Load DICOM | Recalc | Export
 Import from Binary | Import Dose

Status: plan is optimized

Plan

bixel width in [mm]: 5
 Gantry Angle in °: 90 270
 Couch Angle in °: 0 0
 Radiation Mode: protons
 Machine: Generic
 IsoCenter in [mm]: 263.3 265.9 124 Auto.
 # Fractions: 30
 Type of optimization: const_RBExD



3D conformal
 Run Sequencing
 Stratification Levels: 7
 Run Direct Aperture Optimization

Objectives & constraints

+/-	VOI name	VOI type	OP	Function	p	Parameters
-	Rectum	OAR	3	Squared Overdosing	300	d^{max} : 50
-	PTV_68	TARGET	1	Squared Deviation	1000	d^{ref} : 68
-	PTV_56	TARGET	2	Squared Deviation	1000	d^{ref} : 56
-	Bladder	OAR	3	Squared Overdosing	300	d^{max} : 50
-	BODY	OAR	4	Squared Overdosing	100	d^{max} : 30

Visualization

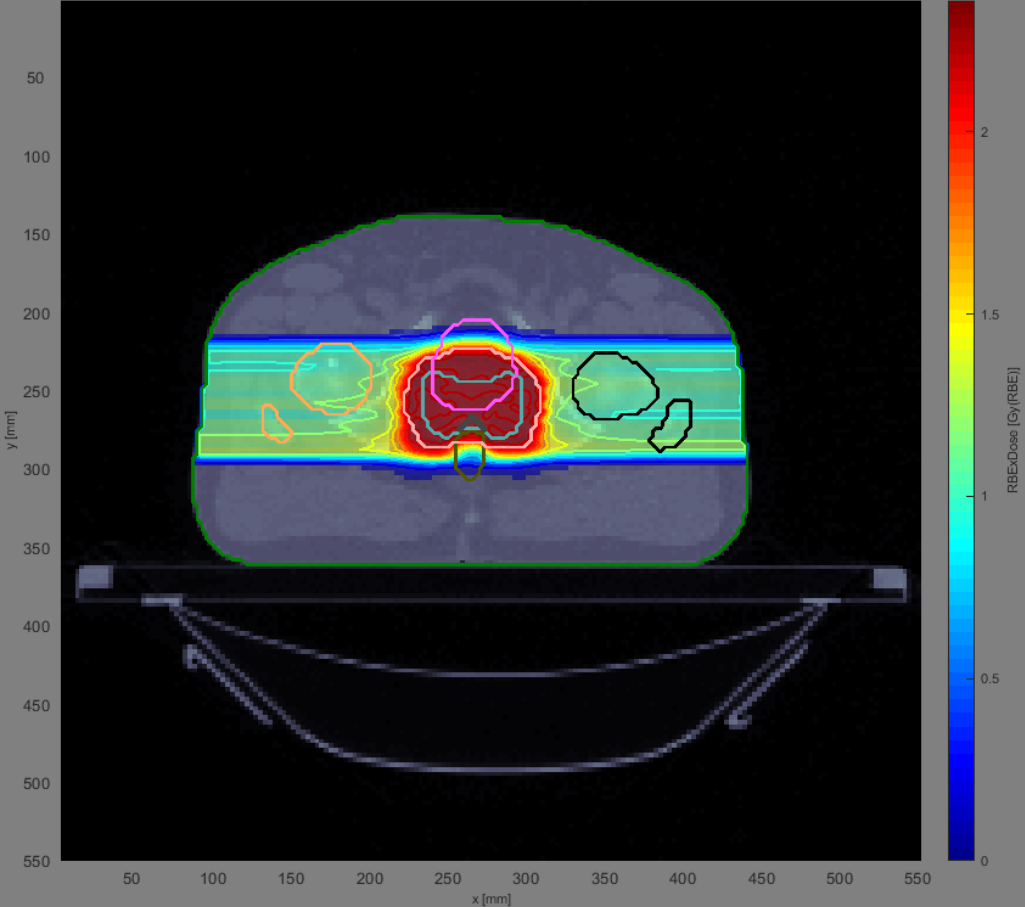
Slice Selection: Type of plot: intensity GoTo: lateral plot CT
 Beam Selection: Plane Selection: axial plot contour
 Offset: Display option: RBExDose plot isolines
 plot dose
 plot isolines labels
 plot iso center
 visualize plan / beams

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Viewing

axial plane z = 99 [mm]



min value: 0
max value: 2.3596

Viewer Options

Result (i.e. dose):
 Window Preset: Custom
 Window Center: 1.18
 Window Width: 2.36
 Range: 0 2.36
 jet
 Lock Settings
 Dose opacity: 1

Structure Visibility

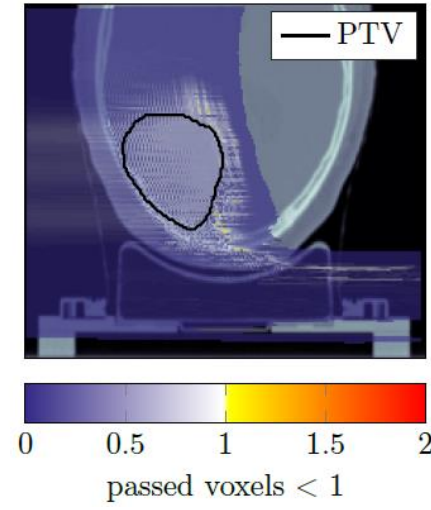
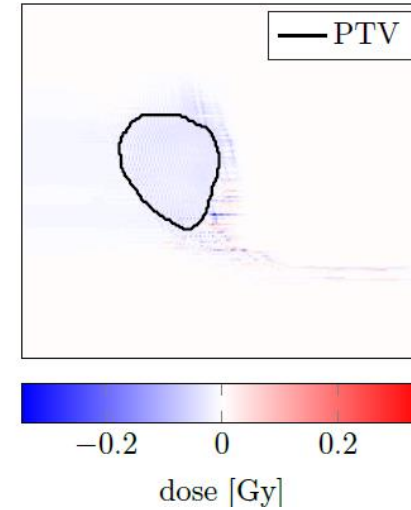
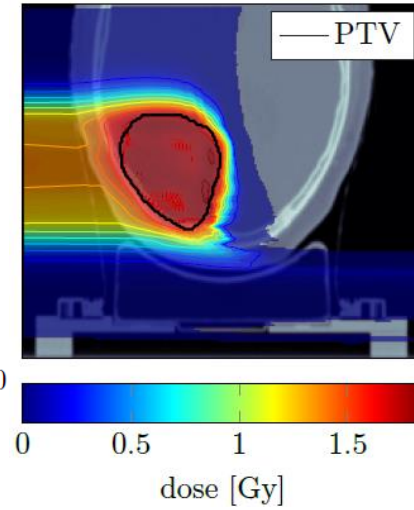
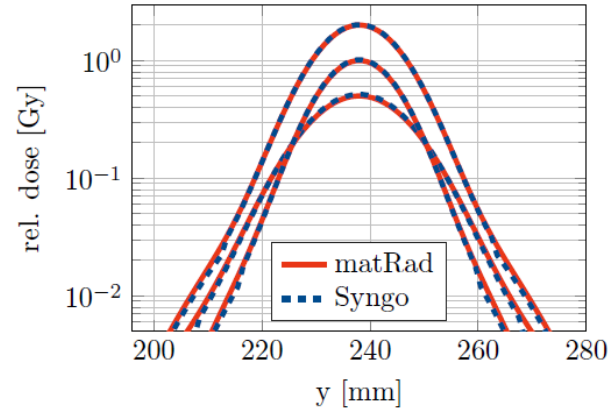
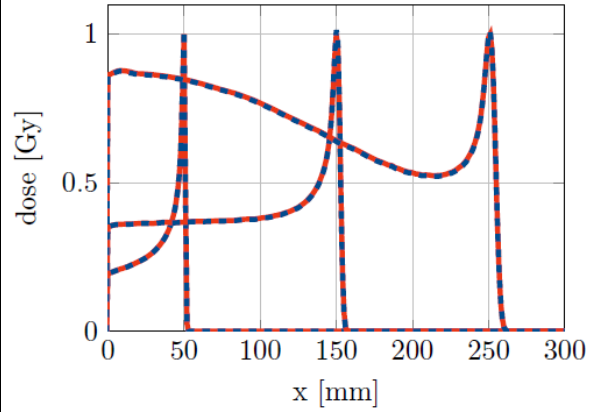
- Rectum
- Penile_bulb
- Lymph Nodes
- Rt femoral head
- prostate_bed
- PTV_68
- PTV_56
- Bladder
- BODY
- Lt femoral head

Info

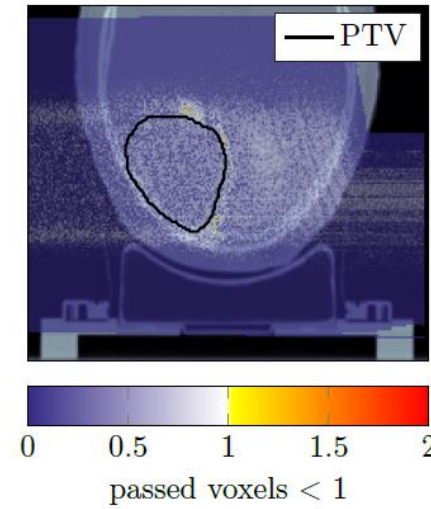
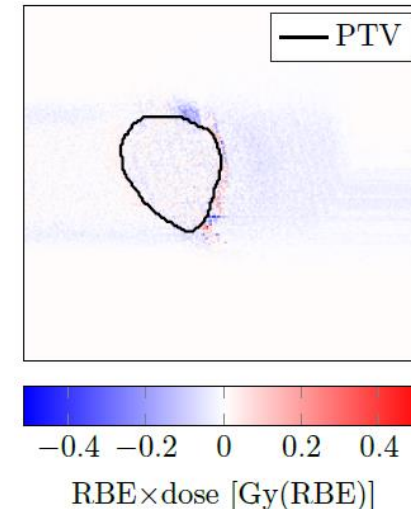
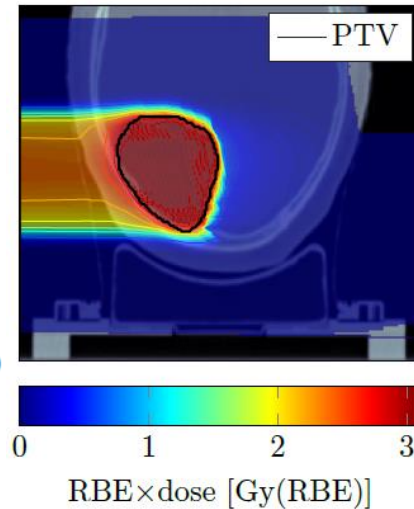
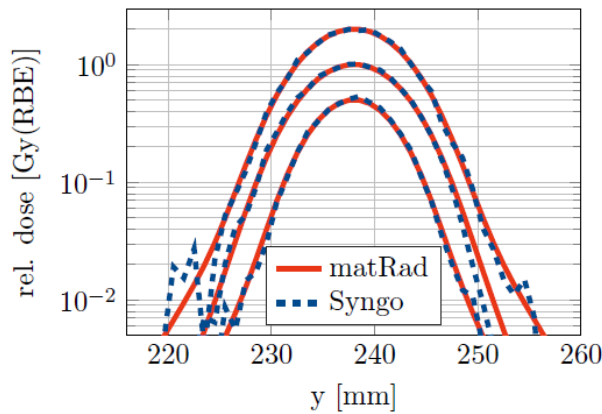
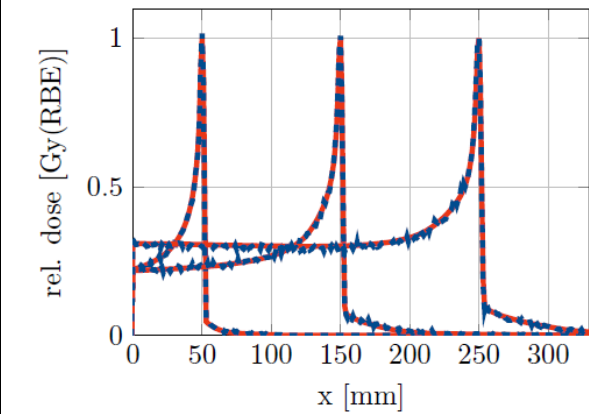
v2.10.0 "Blaise"
(master-c22da7d2)
www.matRad.org

Validation against Syngo Siemens - γ -index > 99.67% (2%,2%)

Protons:



Carbon Ions:



(a) central depth dose profiles

(b) lateral dose profiles

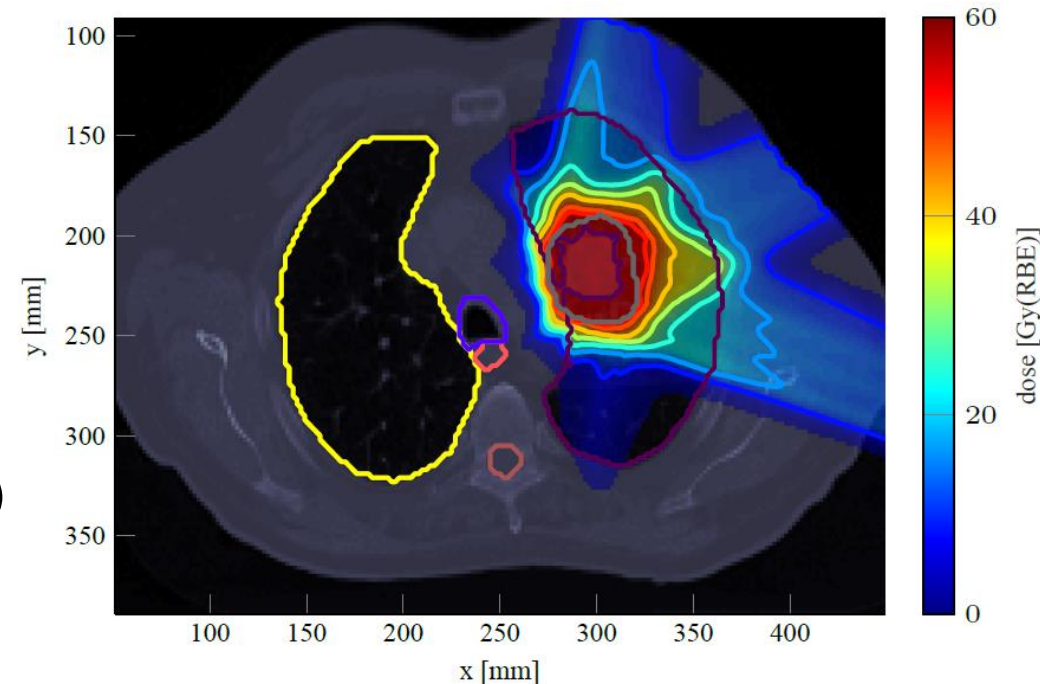
Performance of matRad - Intel Core i7 2.8 GHz, 32 GB RAM

modality	setting	#beams	#bixel	D_{ij} elem. [1e6]	D_{ij} size [GB]	t_{dose} [s]	#iter.	t_{opt} [s]
photons	82 mm D_{ij} samp.	4	2608	172	2.75	295	145	82
photons	40 mm no samp.	4	2608	99	1.59	101	143	44
photons	82 mm D_{ij} samp.	8	3877	426	6.81	741	51	140
photons	40 mm no samp.	8	3877	236	3.77	226	51	66
photons	40 mm D_{ij} samp.	72	13597	567	9.07	853	147	407
protons	99.75 % SG	1	7797	19	0.29	22	123	41
protons	99.75 % DG	1	5955	87	1.38	46	171	109
protons	99.75 % SG	3	28097	56	0.89	68	67	187
protons	99.75 % DG	3	24137	269	4.30	160	262	330
protons	99.75 % SG	2	45574	116	1.86	97	218	137
protons	99.75 % DG	2	27683	520	8.33	299	197	486
carbon	99.75 % SG	1	11780	160	2.55	67	72	92
carbon	99.75 % DG	1	9963	537	8.61	203	79	225
carbon	99.75 % SG	3	42810	411	6.68	310	117	193
carbon	99.75 % DG	3	31205	756	12.1	560	107	365
carbon	99.75 % SG	2	24612	336	5.88	137	177	273
carbon	99.50 % DG	2	16889	855	17.94	472	134	521

Throughput Optimization: 6 GB/s

Development & Research Branches → Hopefully in the next release

- Helium base data (physical & biological)
`-dev_varRBErobOpt`
- Robust / probabilistic optimization & uncertainty quantification
`-dev_varRBErobOpt`
- Variable RBE & effect for protons
`-dev_varRBErobOpt`
- New GUI (Object-oriented & modular, Octave compatible)
`-dev_classGUI`
- Extended MC interfaces (presented last year by Lucas)
`-dev_MonteCarlo`
- External contributions:
 - VMAT `-dev_VMAT(Eric Christiansen)`
 - optimization `-dev_exactOpt (Steven van de Water)`



Proton plan with MCsquare
2e4 histories/beamlet, 4689 beamlets
120 min at (2.5mm)³ resolution

Evaluated on Desktop PC, i7-6700 @ 3.4 GHz (4 cores +HT)

Current Developments at DKFZ

Software Development:

- Unit Testing Framework for Core Functionality
- Python Interface
- Refactoring code for modularization

Future:

- Research Software Engineer position funded by DFG (3 years, WA 4707/1-1)
 - Further professionalization (CI)
 - Helpdesk & Community building
 - ML / AI functionality
 - Interfaces to other open-source software

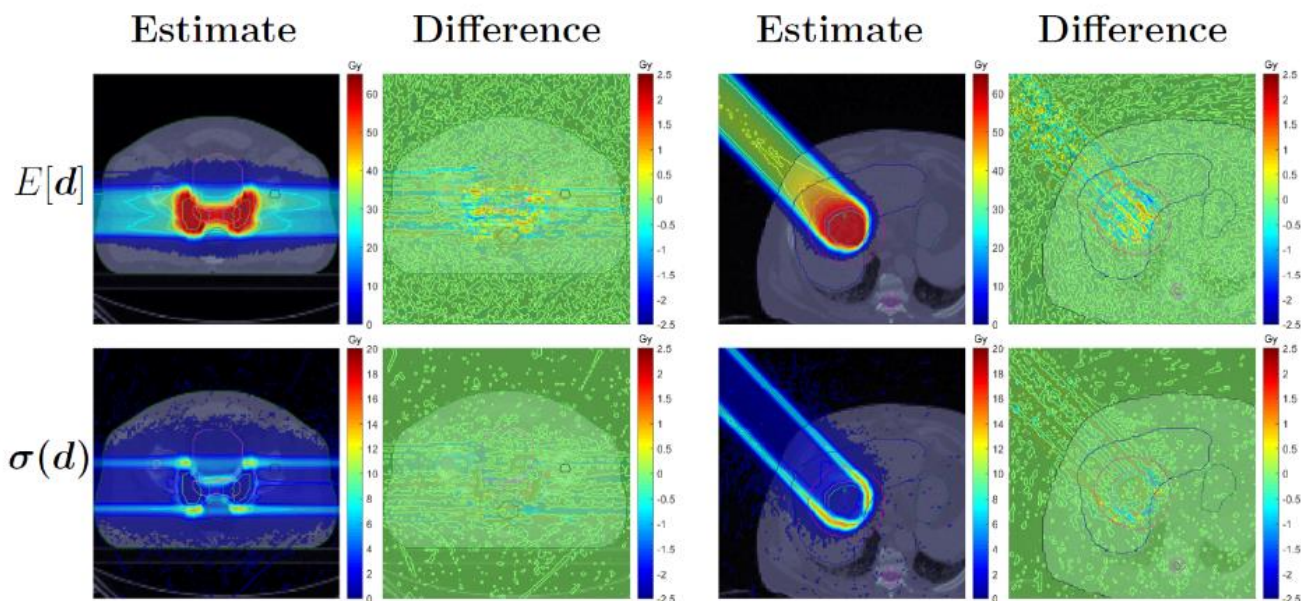
Research-oriented:

- NTCP models & optimization
- Carbon Monte Carlo & Lung degradation
- Joint / mixed-modality / spatio-temporal optimization
 - [Presentation by Amit](#)
- Proton FLASH Planning Tools
- Efficient many-scenario probabilistic planning based on variance objectives
- Superiorization for Inverse Planning
 - [Presentation by Florian](#)

Example matRad Projects

Example Project: MC Dose Uncertainty Quantification (P. Stammer, KIT / DKFZ)

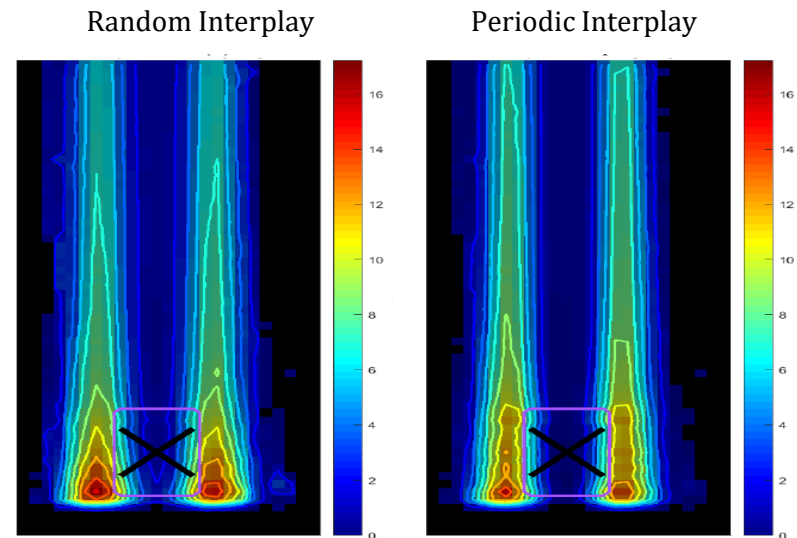
- Avoid computation of explicit error / sample scenarios for robustness analysis and robust optimization with Monte Carlo codes
 → Re-weighting of histories of a single simulation for uncertainty estimation
- Input Uncertainty modeling of static & time-dependent beam application/movement patterns using pencil beam correlations



(a) Prostate

(b) Liver

Estimates of expected dose and variance using the reweighting approach and their difference to a reference [1]



Dose standard deviation in a waterbox for random vs. periodic movement pattern / interplay during treatment

Other Involvements at KIT:

- Development of KiT-RT: A Kinetic Transport Solver for Radiation Therapy (<https://github.com/CSMMLab/KiT-RT>)
- (Dynamical) low rank methods for more time and space efficient UQ in radiative transport

[1] Stammer, P., Burigo, L., Jäkel, O., Frank, M., & Wahl, N. (2021). Efficient uncertainty quantification for Monte Carlo dose calculations using importance (re-) weighting. <https://arxiv.org/abs/2106.11885>

Example Project – Biological dose degradation in lung tissue (N. Homolka)

- Sub-voxel microstructures degradation in Lung in general not captured in deterministic and MC dose calculation

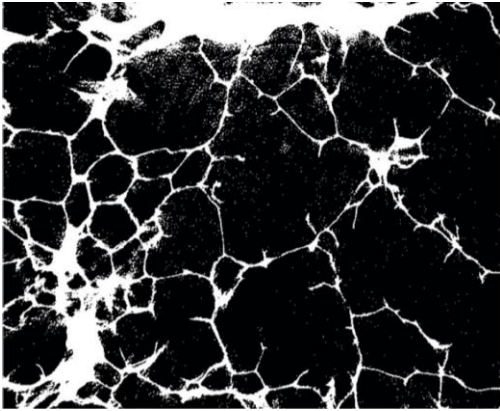


Figure 1: μ CT image of a human lung (binarized) [Baumann et al., 2017].

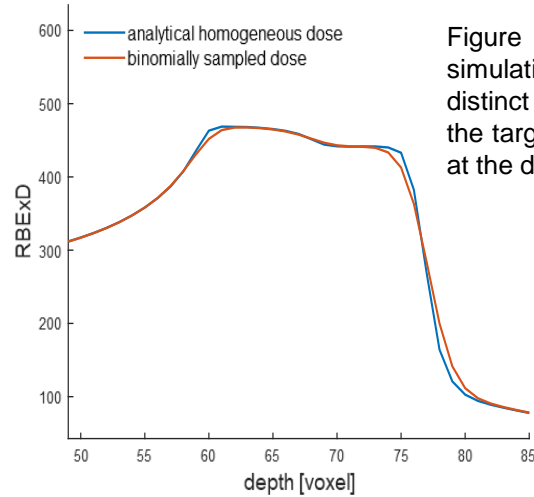


Figure 2: Pencil beam simulations showing the distinct underdosage of the target and overdosage at the distal edge.

- Can be approximated with voxel density & material sampling techniques or analytical convolution techniques

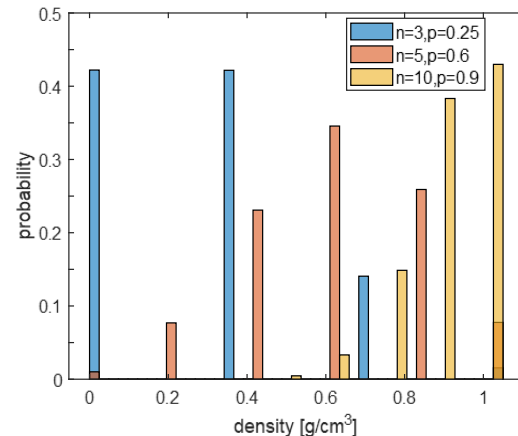


Figure 3: Normalized histogram for 3 different cases for „number of structures in 1 voxel“ and „density of that voxel“

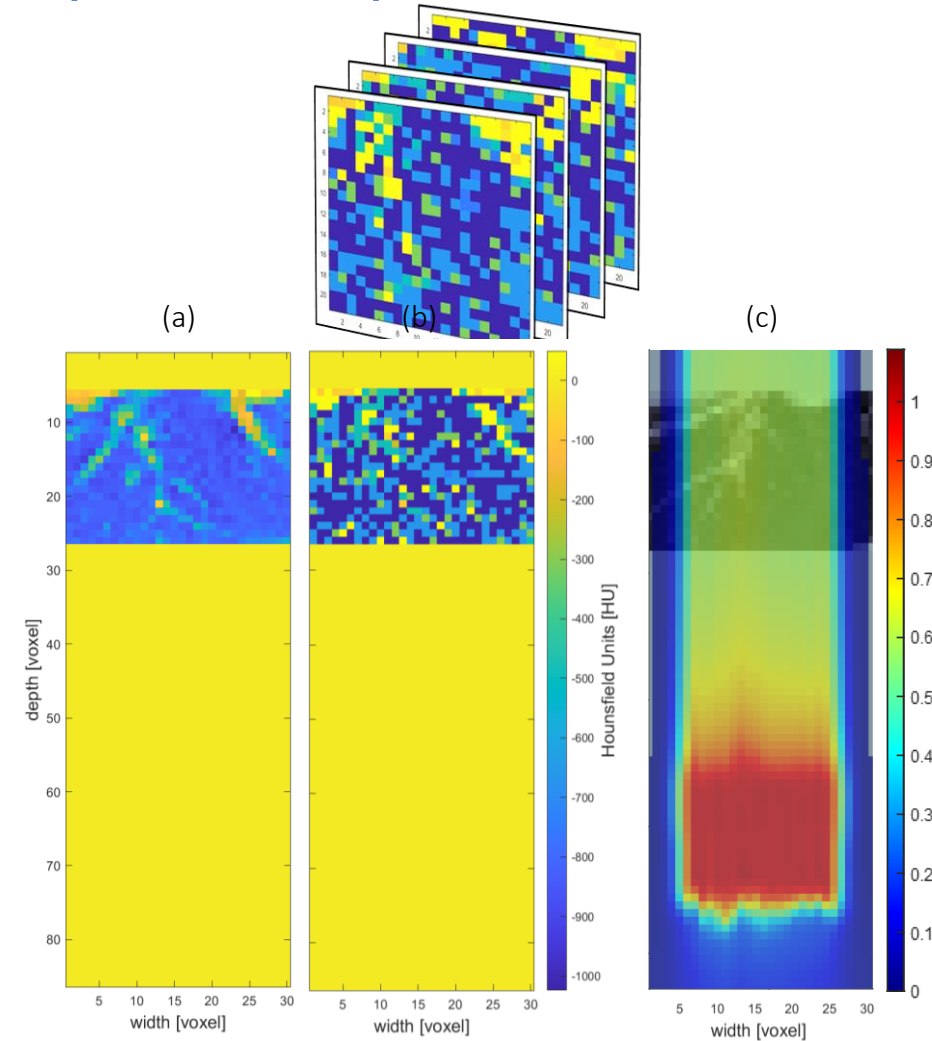


Figure 2: Waterphantom with slab of lung copied from real patient (a), phantom with binomially sampled lung (b), and calculated dose slice averaged over 100 samples (c).

Example Education & Outreach: Particle Therapy Master Class

- Shall become part of the “International Masterclasses – Hands on Particle Physics” (<https://physicsmasterclasses.org/>)
- Educational course for school children
- Introduced by DKFZ/GSI/CERN (Main coordination: Yiota Foka, at DKFZ: Niklas & Joao)
→ First successful stresstests in April 2019 & 2021



Conclusion

- matRad is an open-source treatment planning toolkit with focus on **research & education**
- Used within **internal & external** research projects / collaborations
- **Dose calculation & treatment planning** for photons, protons, helium and carbon (including base data & data import)
- **Active, often research-oriented, development** (internal & external)
 - Monte Carlo interfaces
 - new modalities / optimization techniques
- Efforts in **professionalizing** software development (i. e., continuous integration)

How to get going with matRad?

1. Go to our page on GitHub: www.matRad.org
2. Download the Code, or even better: Familiarize with **git** and Clone
3. Checkout the UI & the code
 - `matRadGUI.m` & `matRad.m`
 - many **examples** in the `examples/` folder
 - Wiki on GitHub: <https://github.com/e0404/matRad/wiki>
4. Ask us from E040-4 for help
5. Profit (and contribute)!



www.matrad.org



matRad

Data IO

DICOM
*.nrrd, *.mha, *.vtk
CERR
VOXELPLAN

Dose calculation

Photons
SVD pencil beam
ompMC interface
Particles
IMPT pencil beam
MCSquare interface
[TOPAS interface](#)
Analytical probabilistic modeling

Analysis & visualization

GUI CT & dose distribution browser
Dose statistics
DVHs

Dose optimization

Fluence and experimental direct aperture optimization
IPOPT <https://projects.coin-or.org/lpopt>
Matlab's proprietary fmincon
Superiorization
Objectives: Quad. dose deviation, mean dose, EUD, DVH
Constraints: Min, max, mean dose, EUD, DVH
Xia, Engel, Siochi MLC sequencer
[Robust and stochastic optimization](#)
[Variable RBE optimization for protons](#)
Coverage based optimization
Analytical probabilistic modeling
VMAT

Base data

Patient data (CT & RTSS)
Photon pencil beam base data
→ <https://github.com/e0404/photonPencilBeamKernelCalc>
Generic proton and carbon ion pencil beam base data
Carbon ion biological base data (LEM IV)
[Helium pencil beam base data](#)
[Helium biological model](#)

Thank you for your attention!

Features in **blue** are available on development branches